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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/940,919

DATE: 09/18/2001

TIME: 10:20:50

Input Set : A:\LEX-0228-USA SEQLIST.txt
Output Set: N:\CRF3\09182001\I940919.raw

4 <110> APPLICANT: Friddle, Carl Johan
 5 Gerhardt, Brenda
 6 Hu, Yi
 8 <120> TITLE OF INVENTION: Novel Human GABA Transporter Protein and Polynucleotides
 Encoding the
 9 Same
 11 <130> FILE REFERENCE: LEX-0228-USA
 C--> 13 <140> CURRENT APPLICATION NUMBER: US/09/940,919 ✓
 C--> 13 <141> CURRENT FILING DATE: 2001-08-28
 13 <150> PRIOR APPLICATION NUMBER: US 60/230,178
 14 <151> PRIOR FILING DATE: 2000-09-01
 16 <160> NUMBER OF SEQ ID NOS: 3
 18 <170> SOFTWARE: FastSEQ for Windows Version 4.0
 20 <210> SEQ ID NO: 1
 21 <211> LENGTH: 1578
 22 <212> TYPE: DNA
 23 <213> ORGANISM: homo sapiens
 25 <400> SEQUENCE: 1
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 27 caggccaaga ttagcgcat ttgcgcagg atgggttttc aggccggccac ggatgaggag 60
 28 gcggtggct tcgcgcattt cgacgaccc tc gactttgagc accggccaggg cctgcagat 120
 29 gacatccctga aagccgaggg agagccctgc ggggacgagg ggcgttgaagc gcccgtcgag 180
 30 ggagacatcc attatcagcg aggcagcgg gtcctctgc cgcctccgg ctccaaggac 240
 31 caggtggag gtggtggcga attcgggggc cacgacaagg ccaaaatcac ggcgtggag 300
 32 gcaggctgga acgtgaccaa cgcgcattccag ggcatttcg tgctgggcct accctacgcc 360
 33 atcctgcacg ggcgttaccc ggggttggcattt ctcatcatct tcgcgcgcgt tgggtgtc 420
 34 tacaccggca agatcctcat cgcgtgcctg tacgaggaga atgaagacgg cgagggtgg 480
 35 cgcgtgcggg actcgatcgat ggccatagcc aacgcctgtc ggcgcgcgcgttccaaacg 540
 36 ctggcgccgc gagtggtgaa cgtgcgcag atcatgcgc tggatgac gtcgcattctg 600
 37 tacgtgggg tgagtggcaa cctcatgtac aacagttcc cggggctgc cgtgtcgac 660
 38 aagtccctgg ccattatcgca caccggccgtg ctgcgcctt ggcgccttc taagaaccc 720
 39 aaggccgtgt ccaaggatcgat tctgcgtgtc actctggccc acttcgtcat caatatctg 780
 40 gtcatagcc actgtctatc gccccgcgc gactggccct ggagaaggta caagttctac 840
 41 atcgacgtca agaaggcccc catctccatt ggcattatcg tggatgac cacttcgtc 900
 42 atcttcctgc ctgcgtggaa gggcaatatcg cagcggccca gcgagttcca ctgcgtatgt 960
 43 aactggacgc acatcgccgc ctgcgtgtc aaggcccttc tgcgcgtcgat cgcctaccc 1020
 44 acctggccgc acgagacccaa ggaggatcgatc acggataacc tggccggcgc catccgcgc 1080
 45 gtggtaaca tctttctggg ggcggccgc ctgttgcctt atcttcgtcc attcttcgtcc 1140
 46 gctgtcgagg tgctggagaa gtcgttc caggaaggca ggcgcgcctt tttccggcc 1200
 47 tgctacagcg ggcacggccgc cctgaagtcc tggggctga cgctgcgtc ggcgcgtcg 1260
 48 gtcctcacgc tgctcatggc catttatgtc cgcgcatttcg cgctgcgtcat gggccctacc 1320
 49 ggcaggccca cggggccgg cctctgttcc ttgtgtccca gccttcata cctgcgcctg 1380
 50 ctctggcgca agctgtgtc gacccaatgc ttcttcgacg tcggccatctt cgtcatcgcc 1440
 51 ggcacatcgca ggcgttccgg ctgcgtgcac tccctcgagg gcctcatcgaa agcctaccga 1500
 52 accaacgcgg aggactag 1560
 54 <210> SEQ ID NO: 2
 55 <211> LENGTH: 525
 56 <212> TYPE: PRT

ENTERED

RAW SEQUENCE LISTING
PATENT APPLICATION:

US/09/940,919

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Input Set : A:\LEX-0228-USA SEQLIST.txt
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57 <213> ORGANISM: homo sapiens
 59 <400> SEQUENCE: 2
 60 Met Ala Thr Leu Leu Arg Ser Lys Leu Ser Asn Val Ala Thr Ser Val
 61 1 5 10 15
 62 Ser Asn Lys Ser Gln Ala Lys Met Ser Gly Met Phe Ala Arg Met Gly
 63 20 25 30
 64 Phe Gln Ala Ala Thr Asp Glu Glu Ala Val Gly Phe Ala His Cys Asp
 65 35 40 45
 66 Asp Leu Asp Phe Glu His Arg Gln Gly Leu Gln Met Asp Ile Leu Lys
 67 50 55 60
 68 Ala Glu Gly Glu Pro Cys Gly Asp Glu Gly Ala Glu Ala Pro Val Glu
 69 65 70 75 80
 70 Gly Asp Ile His Tyr Gln Arg Gly Ser Gly Ala Pro Leu Pro Pro Ser
 71 85 90 95
 72 Gly Ser Lys Asp Gln Val Gly Gly Gly Glu Phe Gly Gly His Asp
 73 100 105 110
 74 Lys Pro Lys Ile Thr Ala Trp Glu Ala Gly Trp Asn Val Thr Asn Ala
 75 115 120 125
 76 Ile Gln Gly Met Phe Val Leu Gly Leu Pro Tyr Ala Ile Leu His Gly
 77 130 135 140
 78 Gly Tyr Leu Gly Leu Phe Leu Ile Ile Phe Ala Ala Val Val Cys Cys
 79 145 150 155 160
 80 Tyr Thr Gly Lys Ile Leu Ile Ala Cys Leu Tyr Glu Glu Asn Glu Asp
 81 165 170 175
 82 Gly Glu Val Val Arg Val Arg Asp Ser Tyr Val Ala Ile Ala Asn Ala
 83 180 185 190
 84 Cys Cys Ala Pro Arg Phe Pro Thr Leu Gly Gly Arg Val Val Asn Val
 85 195 200 205
 86 Ala Gln Ile Ile Glu Leu Val Met Thr Cys Ile Leu Tyr Val Val Val
 87 210 215 220
 88 Ser Gly Asn Leu Met Tyr Asn Ser Phe Pro Gly Leu Pro Val Ser Gln
 89 225 230 235 240
 90 Lys Ser Trp Ser Ile Ile Ala Thr Ala Val Leu Leu Pro Cys Ala Phe
 91 245 250 255
 92 Leu Lys Asn Leu Lys Ala Val Ser Lys Phe Ser Leu Leu Cys Thr Leu
 93 260 265 270
 94 Ala His Phe Val Ile Asn Ile Leu Val Ile Ala Tyr Cys Leu Ser Arg
 95 275 280 285
 96 Ala Arg Asp Trp Ala Trp Glu Lys Val Lys Phe Tyr Ile Asp Val Lys
 97 290 295 300
 98 Lys Phe Pro Ile Ser Ile Gly Ile Ile Val Phe Ser Tyr Thr Ser Gln
 99 305 310 315 320
 100 Ile Phe Leu Pro Ser Leu Glu Gly Asn Met Gln Gln Pro Ser Glu Phe
 101 325 330 335
 102 His Cys Met Met Asn Trp Thr His Ile Ala Ala Cys Val Leu Lys Gly
 103 340 345 350
 104 Leu Phe Ala Leu Val Ala Tyr Leu Thr Trp Ala Asp Glu Thr Lys Glu
 105 355 360 365
 106 Val Ile Thr Asp Asn Leu Pro Gly Ser Ile Arg Ala Val Val Asn Ile

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107	370	375	380
108	Phe Leu Val Ala Lys Ala	Leu Leu Ser Tyr Pro	Leu Pro Phe Phe Ala
109	385	390	395
110	Ala Val Glu Val	Leu Glu Lys Ser	Leu Phe Gln Glu Gly Ser Arg Ala
111		405	400
112	Phe Phe Pro Ala Cys Tyr Ser Gly Asp	Gly Arg Leu Lys Ser Trp Gly	415
113	420	425	430
114	Leu Thr Leu Arg Cys Ala	Leu Val Val Phe Thr Leu Leu Met Ala Ile	
115	435	440	445
116	Tyr Val Pro His Phe Ala	Leu Leu Met Gly Leu Thr GLY Ser Leu Thr	
117	450	455	460
118	Gly Ala Gly Leu Cys Phe	Leu Leu Pro Ser Leu Phe His Leu Arg Leu	
119	465	470	475
120	Leu Trp Arg Lys	Leu Leu Trp His Gln Val Phe Phe Asp Val Ala Ile	480
121	485	490	495
122	Phe Val Ile Gly Gly Ile Cys Ser Val	Ser Gly Phe Val His Ser Leu	
123	500	505	510
124	Glu Gly Leu Ile Glu Ala Tyr Arg Thr Asn Ala Glu Asp		
125	515	520	525
127	<210> SEQ ID NO: 3		
128	<211> LENGTH: 2261		
129	<212> TYPE: DNA		
130	<213> ORGANISM: homo sapiens		
132	<400> SEQUENCE: 3		
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134	cacgggtggg ggcgcgcgc	cagttcgcg tagtgcgtct gacgcgcgtg ccgcgcgcgc	120
135	cgccgcgcgc gcccctcgca	gccccgcgg cagtcgcga gtgcactagc	180
136	caccacgcgc gcccgcgc	ctccgcaga cctgtgcga gttgcccgg tccaggccctg	240
137	agagagcctc gaacgcgc	tgcgagggtc atgagccaga gagccccggg gcgcgcgcgc	300
138	gagagcaagc ggagatagcg	actttgcgc cccagccct cgccttctt catcgcttc	360
139	cccgcatctt cgggtccttc	gtgtccccca cgcgcgcatt ggccaccttgc	420
140	ctccgcagca agctgtccaa	cgtggccacg tccgtgtcca acaagtccca ggccaagatg	480
141	agcggcatgt tcgccaggat	gggtttcag gcggcacagg atgaggaggg ggtgggttgc	540
142	ggcattgcgc acgacccctcgca	ctttgagcac cgccaggggc tgcaatggaa catctgtaaa	600
143	gcccggggc acgcacccctcgca	tttgcggggc gtcggggggc aggctggaa	660
144	tatcagcggag agccctgcgg	ggacgagggc gctgaagcgc ccgtcgaggc agacatccat	720
145	gggtggcaat tcggggggca	tcctctgcgc ccctccggct ccaaggacca ggtggggagg	780
146	gtgaccaacg ccatccagg	aaaatcacgg cgtggggggc aggctggaa	840
147	ggctacctgg ggttgtttct	catgttcgtg ctgggcctac ctaacgcatt cctgcacggc	900
148	atcctcatcg cgtgcgtgt	gcccgcgttgc tgcgtgtcta caccggcaag	960
149	tcgtacgtgg ccatagccaa	cgaggagaat gaagacggcg aggtgggtcg cgtgcgggac	1020
150	gtggtaacg tagcgcagat	cgccctgtgc gccccgcgt tcccaacgcgt gggccggca	1080
151	atgtggcaacc tcatgtacaa	cagttcccg gggctgccc tgtcgcagaa gtccctggcc	1140
152	attatcgcca cggccgtgt	gtgtgcgttgc gccttcattt agaacctcaa ggccgtgtcc	1200
153	aagttcagtc tgctgtgcac	tctggccac ttcgtcatca atatcctggt catagctac	1260
154	tgtctatcgcc gggcgccgcga	ctgggcctgg gagaaggta agttctacat cgacgtcaag	1320
155	aagttcccca tctccattgg	catcatcgta ttcaagtaca cgtctcagat cttcctgcct	1380
156	tcgctggagg gcaatatgca	gcagcccaggc gagttccact gcatgatgaa ctggacgcac	1440
157	atcgcagcct gcgtgctaa	gggccttcc ggcgtcgat cctacccac ctggccgac	1500

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158	gagaccaagg aggtcatcac ggataacctg cccggctcca tccgcgcgt ggtcaacatc	1560
159	tttctgtgg ccaaggcgct gttgtccat cctctgcacat tctttgccgc tgtcgaggtg	1620
160	ctggagaagt cgctctcca ggaaggcagc cgccgccttt tccggcctg ctacagcggc	1680
161	gacgggcgcc tgaagtccctg ggggctgacg ctgcgcgtcg cgctcgctgt cttcacgcgt	1740
162	ctcatggca tttatgtgcc gcacttcgcg ctgcgcattgg gcctcaccgg cagcctcagc	1800
163	ggcgccggcc tctgtttttt gctgcccagc ctcttcacc tggcgcgtcttgcgt ctggcgcaag	1860
164	ctgctgtggc accaagtctt ctgcacgtc gccatcttcg tcacggcg catctgcagc	1920
165	gtgtccggct tcgtgcactc ctcgaggcgtc ctcatcgaaag cctaccgaac caacgcggag	1980
166	gacttagggcg caagggcgag ccccgccgc gcttctgcgc tctctccctt ctccccctcac	2040
167	cccgccccca ccagcccaact ggcgcctgcc ggcgcgcctt ggaggccaag cttaaacat	2100
168	ctctggttcc tagttctga ttattcgggg atggggggga tgggaggggc caggattca	2160
169	cgtccatcg ggcgcgttt tctgtgtcc ttcttttcc acaacaccct ggttttgggg	2220
170	ggaggcgggg tgcatggc ggcagggttc tctgtccttc c	2261

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/940,919

DATE: 09/18/2001
TIME: 10:20:51

Input Set : A:\LEX-0228-USA SEQLIST.txt
Output Set: N:\CRF3\09182001\I940919.raw

L:13 M:270 C: Current Application Number differs, Replaced Current Application No
L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date